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<141> 1999-06-03

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ttc tgc aag tgg cca tgt gag tgc ccg cca tcc cca ccc cgc tgc ccg   95
Phe Cys Lys Trp Pro Cys Glu Cys Pro Pro Ser Pro Pro Arg Cys Pro
                 20                  25                  30

ctg ggg gtc agc ctc atc aca gat ggc tgt gag tgc tgt aag atg tgc   143
Leu Gly Val Ser Leu Ile Thr Asp Gly Cys Glu Cys Cys Lys Met Cys
             35                  40                  45

gct cag cag ctt ggg gac aac tgc acg gag gct gcc atc tgt gac ccc   191
Ala Gln Gln Leu Gly Asp Asn Cys Thr Glu Ala Ala Ile Cys Asp Pro
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cac cgg ggc ctc tac tgt gac tac agc ggg gac cgc ccg agg tac gca   239
His Arg Gly Leu Tyr Cys Asp Tyr Ser Gly Asp Arg Pro Arg Tyr Ala
     65                  70                  75

ata gga gtg tgt gca cag gtg gtc ggt gtg ggc tgc gtc ctg gat ggg   287
Ile Gly Val Cys Ala Gln Val Val Gly Val Gly Cys Val Leu Asp Gly
 80                  85                  90                  95

gtg cgc tac aac aac ggc cag tcc ttc cag cct aac tgc aag tac aac   335
Val Arg Tyr Asn Asn Gly Gln Ser Phe Gln Pro Asn Cys Lys Tyr Asn
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Cys Thr Cys Ile Asp Gly Ala Val Gly Cys Thr Pro Leu Cys Leu Arg
            115                 120                 125

gtg cgc ccc ccg cgt ctc tgg tgc ccc cac ccg cgg cgc gtg agc ata   431
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Pro Arg Lys Thr Ala Pro Arg Asp Thr Gly Ala Phe Asp Ala Val Gly
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gag gtg gag gca tgg cac agg aac tgc ata gcc tac aca agc ccc tgg   575
Glu Val Glu Ala Trp His Arg Asn Cys Ile Ala Tyr Thr Ser Pro Trp
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Ser Pro Cys Ser Thr Ser Cys Gly Leu Gly Val Ser Thr Arg Ile Ser
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ttg cgg cca tgc gat gtg gac atc cat aca ctc att aag gca ggg aag   719
Leu Arg Pro Cys Asp Val Asp Ile His Thr Leu Ile Lys Ala Gly Lys
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aag tgt ctg gct gtg tac cag cca gag gca tcc atg aac ttc aca ctt   767
Lys Cys Leu Ala Val Tyr Gln Pro Glu Ala Ser Met Asn Phe Thr Leu
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gcg ggc tgc atc agc aca cgc tcc tat caa ccc aag tac tgt gga gtt   815
Ala Gly Cys Ile Ser Thr Arg Ser Tyr Gln Pro Lys Tyr Cys Gly Val
                260                 265                 270

tgc atg gac aat agg tgc tgc atc ccc tac aag tct aag act atc gac   863
Cys Met Asp Asn Arg Cys Cys Ile Pro Tyr Lys Ser Lys Thr Ile Asp
            275                 280                 285

gtg tcc ttc cag tgt cct gat ggg ctt ggc ttc tcc cgc cag gtc cta   911
Val Ser Phe Gln Cys Pro Asp Gly Leu Gly Phe Ser Arg Gln Val Leu
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Trp Ile Asn Ala Cys Phe Cys Asn Leu Ser Cys Arg Asn Pro Asn Asp
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atc ttt gct gac ttg gaa tcc tac cct gac ttc tca gaa att gcc aac   1007
Ile Phe Ala Asp Leu Glu Ser Tyr Pro Asp Phe Ser Glu Ile Ala Asn
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         35                  40                  45

Gln Gln Leu Gly Asp Asn Cys Thr Glu Ala Ala Ile Cys Asp Pro His
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Arg Gly Leu Tyr Cys Asp Tyr Ser Gly Asp Arg Pro Arg Tyr Ala Ile
 65                  70                  75                  80

Gly Val Cys Ala Gln Val Val Gly Val Gly Cys Val Leu Asp Gly Val
                 85                  90                  95

Arg Tyr Asn Asn Gly Gln Ser Phe Gln Pro Asn Cys Lys Tyr Asn Cys
            100                 105                 110

Thr Cys Ile Asp Gly Ala Val Gly Cys Thr Pro Leu Cys Leu Arg Val
        115                 120                 125

Arg Pro Pro Arg Leu Trp Cys Pro His Pro Arg Arg Val Ser Ile Pro
    130                 135                 140

Gly His Cys Cys Glu Gln Trp Val Cys Glu Asp Asp Ala Lys Arg Pro
145                 150                 155                 160

Arg Lys Thr Ala Pro Arg Asp Thr Gly Ala Phe Asp Ala Val Gly Glu
                165                 170                 175

Val Glu Ala Trp His Arg Asn Cys Ile Ala Tyr Thr Ser Pro Trp Ser
            180                 185                 190

Pro Cys Ser Thr Ser Cys Gly Leu Gly Val Ser Thr Arg Ile Ser Asn
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Val Asn Ala Gln Cys Trp Pro Glu Gln Glu Ser Arg Leu Cys Asn Leu
    210                 215                 220
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Arg Pro Cys Asp Val Asp Ile His Thr Leu Ile Lys Ala Gly Lys Lys
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Cys Leu Ala Val Tyr Gln Pro Glu Ala Ser Met Asn Phe Thr Leu Ala
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Gly Cys Ile Ser Thr Arg Ser Tyr Gln Pro Lys Tyr Cys Gly Val Cys
            260                 265                 270

Met Asp Asn Arg Cys Cys Ile Pro Tyr Lys Ser Lys Thr Ile Asp Val
        275                 280                 285

Ser Phe Gln Cys Pro Asp Gly Leu Gly Phe Ser Arg Gln Val Leu Trp
    290                 295                 300

Ile Asn Ala Cys Phe Cys Asn Leu Ser Cys Arg Asn Pro Asn Asp Ile
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Phe Ala Asp Leu Glu Ser Tyr Pro Asp Phe Ser Glu Ile Ala Asn
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Thr Phe Thr Pro Ala Pro Leu Glu Glu Thr Thr Thr Arg Pro Glu Phe
        35                  40                  45

Cys Lys Trp Pro Cys Glu Cys Pro Gln Ser Pro Pro Arg Cys Pro Leu
    50                  55                  60

Gly Val Ser Leu Ile Thr Asp Gly Cys Glu Cys Cys Lys Ile Cys Ala
65                  70                  75                  80

Gln Gln Leu Gly Asp Asn Cys Thr Glu Ala Ala Ile Cys Asp Pro His
                85                  90                  95

Arg Gly Leu Tyr Cys Asp Tyr Ser Gly Asp Arg Pro Arg Tyr Ala Ile
            100                 105                 110

Gly Val Cys Ala Gln Val Val Gly Val Gly Cys Val Leu Asp Gly Val
        115                 120                 125

Arg Tyr Thr Asn Gly Glu Ser Phe Gln Pro Asn Cys Arg Tyr Asn Cys
    130                 135                 140

Thr Cys Ile Asp Gly Thr Val Gly Cys Thr Pro Leu Cys Leu Ser Pro
145                 150                 155                 160

Arg Pro Pro Arg Leu Trp Cys Arg Gln Pro Arg His Val Arg Val Pro
                165                 170                 175

Gly Gln Cys Cys Glu Gln Trp Val Cys Asp Asp Asp Ala Arg Arg Pro
            180                 185                 190
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Arg Gln Thr Ala Leu Leu Asp Thr Arg Ala Phe Ala Ala Ser Gly Ala
        195                 200                 205

Val Glu Gln Arg Tyr Glu Asn Cys Ile Ala Tyr Thr Ser Pro Trp Ser
    210                 215                 220

Pro Cys Ser Thr Thr Cys Gly Leu Gly Ile Ser Thr Arg Ile Ser Asn
225                 230                 235                 240

Val Asn Ala Arg Cys Trp Pro Glu Gln Glu Ser Arg Leu Cys Asn Leu
                245                 250                 255

Arg Pro Cys Asp Val Asp Ile Gln Leu His Ile Lys Ala Gly Lys Lys
            260                 265                 270

Cys Leu Ala Val Tyr Gln Pro Glu Glu Ala Thr Asn Phe Thr Leu Ala
        275                 280                 285

Gly Cys Val Ser Thr Arg Thr Tyr Arg Pro Lys Tyr Cys Gly Val Cys
    290                 295                 300

Thr Asp Asn Arg Cys Cys Ile Pro Tyr Lys Ser Lys Thr Ile Ser Val
305                 310                 315                 320

Asp Phe Gln Cys Pro Glu Gly Pro Gly Phe Ser Arg Gln Val Leu Trp
                325                 330                 335

Ile Asn Ala Cys Phe Cys Asn Leu Ser Cys Arg Asn Pro Asn Asp Ile
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Phe Ala Asp Leu Glu Ser Tyr Pro Asp Phe Glu Glu Ile Ala Asn
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Cys Arg Cys Pro Asp Glu Pro Ala Pro Arg Cys Pro Ala Gly Val Ser
        35                  40                  45

Leu Val Leu Asp Gly Cys Gly Cys Cys Arg Val Cys Ala Lys Gln Leu
    50                  55                  60

Gly Glu Leu Cys Thr Glu Arg Asp Pro Cys Asp Pro His Lys Gly Leu
65                  70                  75                  80

Phe Cys Asp Phe Gly Ser Pro Ala Asn Arg Lys Ile Gly Val Cys Thr
                85                  90                  95

Ala Lys Asp Gly Ala Pro Cys Ile Phe Gly Gly Thr Val Tyr Arg Ser
            100                 105                 110

Gly Glu Ser Phe Gln Ser Ser Cys Lys Tyr Gln Cys Thr Cys Leu Asp
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        115                 120                 125

Gly Ala Val Gly Cys Met Pro Leu Cys Ser Met Asp Val Arg Leu Pro
    130                 135                 140

Ser Pro Asp Cys Pro Phe Pro Arg Arg Val Lys Leu Pro Gly Lys Cys
145                 150                 155                 160

Cys Glu Glu Trp Val Cys Asp Glu Pro Lys Asp Gln Thr Val Val Gly
                165                 170                 175

Pro Ala Leu Ala Ala Tyr Arg Leu Glu Asp Thr Phe Gly Pro Asp Pro
            180                 185                 190

Thr Met Ile Arg Ala Asn Cys Leu Val Gln Thr Thr Glu Trp Ser Ala
        195                 200                 205

Cys Ser Lys Thr Cys Gly Met Gly Ile Ser Thr Arg Val Thr Asn Asp
    210                 215                 220

Asn Ala Ser Cys Arg Leu Glu Lys Gln Ser Arg Leu Cys Met Val Arg
225                 230                 235                 240

Pro Cys Glu Ala Asp Leu Glu Glu Asn Ile Lys Lys Gly Lys Lys Cys
                245                 250                 255

Ile Arg Thr Pro Lys Ile Ser Lys Pro Ile Lys Phe Glu Leu Ser Gly
            260                 265                 270

Cys Thr Ser Met Lys Thr Tyr Arg Ala Lys Phe Cys Gly Val Cys Thr
        275                 280                 285

Asp Gly Arg Cys Cys Thr Pro His Arg Thr Thr Thr Leu Pro Val Glu
    290                 295                 300

Phe Lys Cys Pro Asp Gly Glu Val Met Lys Lys Asn Met Met Phe Ile
305                 310                 315                 320

Lys Thr Cys Ala Cys His Tyr Asn Cys Pro Gly Asp Asn Asp Ile Phe
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Glu Ser Leu Tyr Tyr Arg Lys Met Tyr Gly Asp Met Ala
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Pro Leu Glu Ala Pro Lys Cys Ala Pro Gly Val Gly Leu Val Arg Asp
         35                  40                  45

Gly Cys Gly Cys Cys Lys Val Cys Ala Lys Gln Leu Asn Glu Asp Cys
     50                  55                  60
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Ser Lys Thr Gln Pro Cys Asp His Thr Lys Gly Leu Glu Cys Asn Phe
 65                  70                  75                  80

Gly Ala Ser Ser Thr Ala Leu Lys Gly Ile Cys Arg Ala Gln Ser Glu
                85                  90                  95

Gly Arg Pro Cys Glu Tyr Asn Ser Arg Ile Tyr Gln Asn Gly Glu Ser
            100                 105                 110

Phe Gln Pro Asn Cys Lys His Gln Cys Thr Cys Ile Asp Gly Ala Val
        115                 120                 125

Gly Cys Ile Pro Leu Cys Pro Gln Glu Leu Ser Leu Pro Asn Leu Gly
    130                 135                 140

Cys Pro Asn Pro Arg Leu Val Lys Val Thr Gly Gln Cys Cys Glu Glu
145                 150                 155                 160

Trp Val Cys Asp Glu Asp Ser Ile Lys Asp Pro Met Glu Asp Gln Asp
                165                 170                 175

Gly Leu Leu Gly Lys Glu Leu Gly Phe Asp Ala Ser Glu Val Glu Leu
            180                 185                 190

Thr Arg Asn Asn Glu Leu Ile Ala Val Gly Lys Gly Ser Ser Leu Lys
        195                 200                 205

Arg Leu Pro Val Phe Gly Met Glu Pro Arg Ile Leu Tyr Asn Pro Leu
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Gln Gly Gln Lys Cys Ile Val Gln Thr Thr Ser Trp Ser Gln Cys Ser
225                 230                 235                 240

Lys Thr Cys Gly Thr Gly Ile Ser Thr Arg Val Thr Asn Asp Asn Pro
                245                 250                 255

Glu Cys Arg Leu Val Lys Glu Thr Arg Ile Cys Glu Val Arg Pro Cys
            260                 265                 270

Gly Gln Pro Val Tyr Ser Ser Leu Lys Lys Gly Lys Lys Cys Ser Lys
        275                 280                 285

Thr Lys Lys Ser Pro Glu Pro Val Arg Phe Thr Tyr Ala Gly Cys Leu
    290                 295                 300

Ser Val Lys Lys Tyr Arg Pro Lys Tyr Cys Gly Ser Cys Val Asp Gly
305                 310                 315                 320

Arg Cys Cys Thr Pro Gln Leu Thr Arg Thr Val Lys Met Arg Phe Arg
                325                 330                 335

Cys Glu Asp Gly Glu Thr Phe Ser Lys Asn Val Met Met Ile Gln Ser
            340                 345                 350

Cys Lys Cys Asn Tyr Asn Cys Pro His Ala Asn Glu Ala Ala Phe Pro
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             20                  25                  30

Gln Arg Cys Pro Pro Gln Cys Pro Gly Arg Cys Pro Ala Thr Pro Pro
         35                  40                  45

Thr Cys Ala Pro Gly Val Arg Ala Val Leu Asp Gly Cys Ser Cys Cys
     50                  55                  60

Leu Val Cys Ala Arg Gln Arg Gly Glu Ser Cys Ser Asp Leu Glu Pro
 65                  70                  75                  80

Cys Asp Glu Ser Ser Gly Leu Tyr Cys Asp Arg Ser Ala Asp Pro Ser
                 85                  90                  95

Asn Gln Thr Gly Ile Cys Thr Ala Val Glu Gly Asp Asn Cys Val Phe
            100                 105                 110

Asp Gly Val Ile Tyr Arg Ser Gly Glu Lys Phe Gln Pro Ser Cys Lys
        115                 120                 125

Phe Gln Cys Thr Cys Arg Asp Gly Gln Ile Gly Cys Val Pro Arg Cys
    130                 135                 140

Gln Leu Asp Val Leu Leu Pro Glu Pro Asn Cys Pro Ala Pro Arg Lys
145                 150                 155                 160

Val Glu Val Pro Gly Glu Cys Cys Glu Lys Trp Ile Cys Gly Pro Asp
                165                 170                 175

Glu Glu Asp Ser Leu Gly Gly Leu Thr Leu Ala Ala Tyr Arg Pro Glu
            180                 185                 190

Ala Thr Leu Gly Val Glu Val Ser Asp Ser Ser Val Asn Cys Ile Glu
        195                 200                 205

Gln Thr Thr Glu Trp Thr Ala Cys Ser Lys Ser Cys Gly Met Gly Phe
    210                 215                 220

Ser Thr Arg Val Thr Asn Arg Asn Arg Gln Cys Glu Met Leu Lys Gln
225                 230                 235                 240

Thr Arg Leu Cys Met Val Arg Pro Cys Glu Gln Glu Pro Glu Gln Pro
                245                 250                 255

Thr Asp Lys Lys Gly Lys Lys Cys Leu Arg Thr Lys Lys Ser Leu Lys
            260                 265                 270

Ala Ile His Leu Gln Phe Lys Asn Cys Thr Ser Leu His Thr Tyr Lys
        275                 280                 285

Pro Arg Phe Cys Gly Val Cys Ser Asp Gly Arg Cys Cys Thr Pro His
    290                 295                 300

Asn Thr Lys Thr Ile Gln Ala Glu Phe Gln Cys Ser Pro Gly Gln Ile
305                 310                 315                 320
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Val Lys Lys Pro Val Met Val Ile Gly Thr Cys Thr Cys His Thr Asn
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<221> misc_feature
<222> (483)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (488)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (491)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (494)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (502)
<223> n equals a, t, g or c

<400> 8
ggcanagttt ttttttcca ttatatttat tattcttaat ttttattgaa tcaggccaca 60

ccattgcagg gaaattctga cttagaaaga gcttcattac attggatcca aatttccttc 120

tccatggctt caacccatgg gtctgagact tgccctccgg ntnctgcata agatagttcc 180

gtttctcaaa accaatggcg actgggttat tctgaccacc ttcctccacc cattgccaga 240

ataatcccta gtttctccaa ccagacttcc tatgatcttt gaaagctaag ttcatttccc 300

aattgagatg caattccagt aagaaccaag ccttggggtt nccanggatt tcaatgggnt 360

gngcgttnnc cagcntgnaa ttggnaaagg caggggtttn caccccggga aaaccaaaag 420

ggttccaccg gcttnacgag gggccntcca gggggaatnc ctttaanaag atctggaagg 480

gancccantt ngtnaaaaaa gncttctgga aaaaagcctt gcaggctaaa aatgggggg 539

<210> 9

<211> 311
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (177)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (189)
<223> n equals a, t, g or c

<400> 9
attgaatgtt gtgtagttat tcacagggaa ttctgtgcag tgtgcagaga gattcctaaa 60

cgggaaaagg actgggaata catcctcctt actgtgacct ccccaaaacc tagtccagtg 120

caaggtatac agtggtgctc attaaatact tgatgaatac aggaagctgt gcatgtnttc 180

ctacttttnt tcgaagctct cttcttccaa agctacatga aaatagaatt ttaacagtca 240

aaattttata ttaagtgcct tagcaaaaga gacatttaat attttcaaag aaatgcatat 300

gtatgtatac a 311

<210> 10
<211> 197
<212> DNA
<213> Homo sapiens

<400> 10
ctcttctgta agtcagtgtg aatcatgtta gattttctga gagtgaaaac acctgccatc 60

tacaaattac aaggctggat aacagctcac tccatttgaa attcagtgga aacccaagag 120

ctaggttctt actggaattt gcatctcaat ttgggaaact gaacttagct ttcaaagatc 180

ataggaagtc ttgttgg 197

<210> 11
<211> 484
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (39)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (75)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (88)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (104)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (111)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (145)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (165)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (170)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (229)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (239)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (239)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (328)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (337)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (356)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (377)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (379)

<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (392)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (397)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (418)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (424)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (476)
<223> n equals a, t, g or c

<400> 11
ggcagagtgt atacatatat ttgtgtatgc gtatgaagna attcttgtat aaagagaatt 60

cactccatga atganctctt ctgtagtnna gtgtgaatca tgtnagattt nctaagagtg 120

aaaaacacct gccatctaca aattnacaag gctggataac agctncactn ccatttgaaa 180

attcagtggg aaacccaaga gctaggttct tactggaatt tgccatctnc aatttgggna 240

aactgaaact taggctttcc aaaggttcat agggaagtct gggttggagg aaactagggg 300

attattcctg ggcaatgggg tgggaggnag gtgggtncag aattaacccc gttcgncctt 360

tggttttgag gaacggnant atcttatggc gngcccnggg gaagttcttc ggaccctngg 420

gttnnaggcc tgggggaggg aattttgggt cccatgtatg aggtctttct aggtcnggat 480

ttcc 484

<210> 12
<211> 236
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (38)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (182)
<223> n equals a, t, g or c

<220>

<221> misc_feature
<222> (216)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (220)
<223> n equals a, t, g or c

<220>
<221> misc_feature
<222> (228)
<223> n equals a, t, g or c

<400> 12
ggcacagcgc tcagcagctt ggggacaact gcacggangc tgccatctgt gacccccacc 60

ggccgctgct actgtgacta catcggggac ccacgaggta cgcaataggg agtgtgtgca 120

caggtggtcg gtgtgggctg cgtcctggga tggggtgcgc tacaacaacg gaccagtcct 180

tnccagccta aactggcaat gacaactgcc acgtgncatn cggacggnna cggtgg 236

<210> 13
<211> 32
<212> DNA
<213> Homo sapiens

<400> 13
cgcggatccg cgatggactt taccccagct cc 32

<210> 14
<211> 39
<212> DNA
<213> Homo sapiens

<400> 14
ctagtctaga ctaggttggc aatttctgag aagtcaggg 39

<210> 15
<211> 30
<212> DNA
<213> Homo sapiens

<400> 15
cgcggatccg cgcgacttta ccccagctcc 30

<210> 16
<211> 39
<212> DNA
<213> Homo sapiens

<400> 16
ctagggtacc ctaggttggc aatttctgag aagtcaggg 39

<210> 17
<211> 733
<212> DNA

<213> Homo sapiens

<400> 17
gggatccgga gcccaaatct tctgacaaaa ctcacacatg cccaccgtgc ccagcacctg 60

aattcgaggg tgcaccgtca gtcttcctct tccccccaaa acccaaggac accctcatga 120

tctcccggac tcctgaggtc acatgcgtgg tggtggacgt aagccacgaa gaccctgagg 180

tcaagttcaa ctggtacgtg gacggcgtgg aggtgcataa tgccaagaca aagccgcggg 240

aggagcagta caacagcacg taccgtgtgg tcagcgtcct caccgtcctg caccaggact 300

ggctgaatgg caaggagtac aagtgcaagg tctccaacaa agccctccca acccccatcg 360

agaaaaccat ctccaaagcc aaagggcagc cccgagaacc acaggtgtac accctgcccc 420

catcccggga tgagctgacc aagaaccagg tcagcctgac ctgcctggtc aaaggcttct 480

atccaagcga catcgccgtg gagtgggaga gcaatgggca gccggagaac aactacaaga 540

ccacgcctcc cgtgctggac tccgacggct ccttcttcct ctacagcaag ctcaccgtgg 600

acaagagcag gtggcagcag gggaacgtct tctcatgctc cgtgatgcat gaggctctgc 660

acaaccacta cacgcagaag agcctctccc tgtctccggg taaatgagtg cgacggccgc 720

gactctagag gat 733

<210> 18
<211> 5
<212> PRT
<213> Homo sapiens

<220>
<221> SITE
<222> (3)
<223> Xaa equals any amino acid

<400> 18
Trp Ser Xaa Trp Ser
1 5

<210> 19
<211> 86
<212> DNA
<213> Homo sapiens

<400> 19
gcgcctcgag atttccccga aatctagatt tccccgaaat gatttccccg aaatgatttc 60

cccgaaatat ctgccatctc aattag 86

<210> 20
<211> 27
<212> DNA
<213> Homo sapiens

<400> 20
gcggcaagct ttttgcaaag cctaggc 27

<210> 21
<211> 271
<212> DNA
<213> Homo sapiens

<400> 21
ctcgagattt ccccgaaatc tagatttccc cgaaatgatt tccccgaaat gatttccccg 60

aaatatctgc catctcaatt agtcagcaac catagtcccg cccctaactc cgcccatccc 120

gcccctaact ccgcccagtt ccgcccattc tccgccccat ggctgactaa ttttttttat 180

ttatgcagag gccgaggccg cctcggcctc tgagctattc cagaagtagt gaggaggctt 240

ttttggaggc ctaggctttt gcaaaaagct t 271

<210> 22
<211> 32
<212> DNA
<213> Homo sapiens

<400> 22
gcgctcgagg gatgacagcg atagaacccc gg 32

<210> 23
<211> 31
<212> DNA
<213> Homo sapiens

<400> 23
gcgaagcttc gcgactcccc ggatccgcct c 31

<210> 24
<211> 12
<212> DNA
<213> Homo sapiens

<400> 24
ggggactttc cc 12

<210> 25
<211> 73
<212> DNA
<213> Homo sapiens

<400> 25
gcggcctcga ggggactttc ccggggactt tccggggact ttccgggact ttccatcctg 60

ccatctcaat tag 73

<210> 26
<211> 27
<212> DNA
<213> Homo sapiens

<400> 26

gcggcaagct ttttgcaaag cctaggc 27

<210> 27
<211> 256
<212> DNA
<213> Homo sapiens

<400> 27
ctcgagggga ctttcccggg gactttccgg ggactttccg ggactttcca tctgccatct 60

caattagtca gcaaccatag tcccgccccct aactccgccc atcccgcccc taactccgcc 120

cagttccgcc cattctccgc cccatggctg actaattttt tttatttatg cagaggccga 180

ggccgcctcg gcctctgagc tattccagaa gtagtgagga ggcttttttg gaggcctagg 240

cttttgcaaa aagctt 256